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What is claimed is:

- 1. A method for the detection of one or more NF-kB regulatory factors comprising the steps of:
 - a) providing a slimb protein, and a sample suspected of containing one or more NF-κB regulatory factors; and
 - b) exposing said slimb protein to said sample under conditions such that said slimb protein binds to said one or more NF-κB regulatory factors to form a slimb/regulatory factor complex.
- 2. The method of Claim 1, further comprising the step of detecting said slimb/regulatory factor complex.
- 3. The method of Claim 1, further comprising the step of observing said slimb/regulatory factor complex for degradation of said one or more NF-kB regulatory factors.
- 4. The method of Claim 1, further comprising the step of exposing said slimb protein and one or more NF-κB regulatory factors to an F-box protein antagonist.
- 5. The method of Claim 4, wherein said F-box protein antagonist prevents the formation of said slimb/regulatory factor complex.

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A method for the detection of a slimb protein complex, comprising the

- providing a slimb protein and a sample suspected of containing one or a) more proteins capable of forming a complex with said slimb protein; and
- exposing said slimb protein to said one or more proteins capable of b) forming a complex with\said slimb protein under conditions such that said slimb protein binds to said one or more proteins capable of forming a complex with said slimb\protein to form a slimb protein complex.
- The method of Claim 6, further comprising the step of detecting said 7. slimb protein complex.
- The method of Claim 6, wherein step b) further comprises exposing said 8. slimb protein and said one or there proteins capable of forming a complex with said slimb protein to an F-box protein antagonist.
- The method of Claim 8, wherein said F-box protein antagonist prevents 9. the formation of said slimb protein complex.
- An isolated nucleotide sequence comprising nucleotide sequence 10. encoding at least one functionally active fragement of an F-box protein, wherein said sequence consists of a least a portion of the sequence set forth in SEQ ID NOS: 54 and 56.

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